
Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2009; month=11; day=25; hr=11; min=14; sec=7; ms=427;]

Validated By CRFValidator v 1.0.3

Application No: 10598965 Version No: 4.0

Input Set:

Output Set:

Started: 2009-11-24 16:47:07.748

Finished: 2009-11-24 16:47:10.456

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 708 ms

Total Warnings: 31

Total Errors: 1

No. of SeqIDs Defined: 32

Actual SeqID Count: 32

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W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(4)
M	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(5)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(6)
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W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(8)
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W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(12)
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W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(19)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(20)
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Input Set:

Output Set:

Started: 2009-11-24 16:47:07.748

Finished: 2009-11-24 16:47:10.456

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 708 ms

Total Warnings: 31

Total Errors: 1

No. of SeqIDs Defined: 32

Actual SeqID Count: 32

Error code Error Description

This error has occured more than 20 times, will not be displayed

E 257 Invalid sequence data feature in <221> in SEQ ID (32)

SEQUENCE LISTING

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Lys	Thr	Tyr	Val 20	Asp	Ser	Leu	Asn	Val 25	Ile	Arg	Ser	Ala	Ile 30	Gly	Thr
Pro	Leu	Gln 35	Thr	Ile	Ser	Ser	Gly 40	Gly	Thr	Ser	Leu	Leu 45	Met	Ile	Asp
Ser	Gly 50	Ser	Gly	Asp	Asn	Leu 55	Phe	Ala	Val	Asp	Val 60	Arg	Gly	Ile	Asp
Pro 65	Glu	Glu	Gly	Arg	Phe 70	Asn	Asn	Leu	Arg	Leu 75	Ile	Val	Glu	Arg	Asn
Asn	Leu	Tyr	Val	Thr 85	Gly	Phe	Val	Asn	Arg 90	Thr	Asn	Asn	Val	Phe 95	Tyr
Arg	Phe	Ala	Asp 100	Phe	Ser	His	Val	Thr 105	Phe	Pro	Gly	Thr	Thr 110	Ala	Val

Thr Leu Ser Gly Asp Ser Ser Tyr Thr Thr Leu Gln Arg Val Ala Gly

125

120

115

Ile Ser Arg Thr Gly Met Gln Ile Asn Arg His Ser Leu Thr Thr Ser 130 135 140

Tyr Leu Asp Leu Met Ser His Ser Gly Thr Ser Leu Thr Gln Ser Val
145 150 150 160

Ala Arg Ala Met Leu Arg Phe Val Thr Val Thr Ala Glu Ala Leu Arg
165 170 175

Phe Arg Gln Ile Gln Arg Gly Phe Arg Thr Thr Leu Asp Asp Leu Ser 180 185 190

Gly Arg Ser Tyr Val Met Thr Ala Glu Asp Val Asp Leu Thr Leu Asn 195 200 205

Trp Gly Arg Leu Ser Ser Val Leu Pro Asp Tyr His Gly Gln Asp Ser 210 220

Val Arg Val Gly Arg Ile Ser Phe Gly Ser Ile Asn Ala Ile Leu Gly 225 230 230 235 235 240

Ser Val Ala Leu Ile Leu Asn Cys His His His Ala Ser Arg Val Ala 245 250 255

Arg Met Ala Ser Asp Glu Phe Pro Ser Met Cys Pro Ala Asp Gly Arg 260 270

Val Arg Gly Ile Thr His Asn Lys Ile Leu Trp Asp Ser Ser Thr Leu 275 280 285

Gly Ala Ile Leu Met Arg Arg Thr Ile Ser Ser 290 295

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<223> Description of Artificial Sequence: Synthetic primer

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                                     10
                                                         15
1
Arg Ala Ser Lys Glu Phe Thr Leu Asp Phe Ser Thr Ala Lys Thr Tyr
            20
                                25
                                                     30
Val Asp Ser Leu Asn Val Ile Arg Ser Ala Ile Gly Thr Pro Leu Gln
                            40
        35
                                                 45
Thr Ile Ser Ser Gly Gly Thr Ser Leu Leu Met Ile Asp Ser Gly Ser
    50
                        55
                                             60
Gly Asp Asn Leu Phe Ala Val Asp Val Arg Gly Ile Asp Pro Glu Glu
65
                                         75
                                                             80
                    70
Gly Arg Phe Asn Asn Leu Arg Leu Ile Val Glu Arg Asn Asn Leu Tyr
                85
                                     90
                                                         95
Val Thr Gly Phe Val Asn Arg Thr Asn Asn Val Phe Tyr Arg Phe Ala
            100
                                105
                                                     110
Asp Phe Ser His Val Thr Phe Pro Gly Thr Thr Ala Val Thr Leu Ser
        115
                            120
                                                 125
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Gly Asp Ser Ser Tyr Thr Thr Leu Gln Arg Val Ala Gly Ile Ser Arg

31

130 135 140

Thr Gly Met Gln Ile Asn Arg His Ser Leu Thr Thr Ser Tyr Leu Asp 145 150 155 160

Leu Met Ser His Ser Gly Thr Ser Leu Thr Gln Ser Val Ala Arg Ala

165
170
175

Met Leu Arg Phe Val Thr Val Thr Ala Glu Ala Leu Arg Phe Arg Gln
180 185 190

Ile Gln Arg Gly Phe Arg Thr Thr Leu Asp Asp Leu Ser Gly Arg Ser 195 200 205

Tyr Val Met Thr Ala Glu Asp Val Asp Leu Thr Leu Asn Trp Gly Arg 210 215 220

Leu Ser Ser Val Leu Pro Asp Tyr His Gly Gln Asp Ser Val Arg Val 225 230 230 235 240

Gly Arg Ile Ser Phe Gly Ser Ile Asn Ala Ile Leu Gly Ser Val Ala
245 250 255

Leu Ile Leu Asn Cys His His His Ile Tyr Ser Asn Lys Leu Met Ala 260 270

Ser Arg Val Ala Arg Met Ala Ser Asp Glu Phe Pro Ser Met Cys Pro 275 280 285

Ala Asp Gly Arg Val Arg Gly Ile Thr His Asn Lys Ile Leu 290 295 300

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<211> 319

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Val	Asp	Ser 35	Leu	Asn	Val	Ile	Arg 40	Ser	Ala	Ile	Gly	Thr 45	Pro	Leu	Gln
Thr	Ile 50	Ser	Ser	Gly	Gly	Thr 55	Ser	Leu	Leu	Met	Ile 60	Asp	Ser	Gly	Ser
Gly 65	Asp	Asn	Leu	Phe	Ala 70	Val	Asp	Val	Arg	Gly 75	Ile	Asp	Pro	Glu	Glu 80
Gly	Arg	Phe	Asn	Asn 85	Leu	Arg	Leu	Ile	Val 90	Glu	Arg	Asn	Asn	Leu 95	Tyr
Val	Thr	Gly	Phe 100	Val	Asn	Arg	Thr	Asn 105	Asn	Val	Phe	Tyr	Arg 110	Phe	Ala
Asp	Phe	Ser 115	His	Val	Thr	Phe	Pro 120	Gly	Thr	Thr	Ala	Val 125	Thr	Leu	Ser
Gly	Asp 130	Ser	Ser	Tyr	Thr	Thr 135	Leu	Gln	Arg	Val	Ala 140	Gly	Ile	Ser	Arg
Thr 145	Gly	Met	Gln	Ile	Asn 150	Arg	His	Ser	Leu	Thr 155	Thr	Ser	Tyr	Leu	Asp 160
Leu	Met	Ser	His	Ser 165	Gly	Thr	Ser	Leu	Thr 170	Gln	Ser	Val	Ala	Arg 175	Ala
Met	Leu	Arg	Phe 180	Val	Thr	Val	Thr	Ala 185	Glu	Ala	Leu	Arg	Phe 190	Arg	Gln
Ile	Gln	Arg 195	Gly	Phe	Arg	Thr	Thr 200	Leu	Asp	Asp	Leu	Ser 205	Gly	Arg	Ser
Tyr	Val 210	Met	Thr	Ala	Glu	Asp 215	Val	Asp	Leu	Thr	Leu 220	Asn	Trp	Gly	Arg
Leu 225	Ser	Ser	Val	Leu	Pro 230	Asp	Tyr	His	Gly	Gln 235	Asp	Ser	Val	Arg	Val 240

Gly Arg Ile Ser Phe Gly Ser Ile Asn Ala Ile Leu Gly Ser Val Ala

245 250 255

Leu Ile Leu Asn Cys His His Ala Ala Phe Ala Asp Leu Ile Ala 265 260 270 Ser Arg Val Ala Arg Met Ala Ser Asp Glu Phe Pro Ser Met Cys Pro 275 280 285 Ala Asp Gly Arg Val Arg Gly Ile Thr His Asn Lys Ile Leu Trp Asp 290 295 300 Ser Ser Thr Leu Gly Ala Ile Leu Met Arg Arg Thr Ile Ser Ser 305 310 315 <210> 6 <211> 7 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic peptide <400> 6 Ile Tyr Ser Asn Lys Leu Met 1 5 <210> 7 <211> 7 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic peptide <400> 7 Ala Ala Phe Ala Asp Leu Ile 1 <210> 8 <211> 7 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic peptide

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Cys His His Pro Asp Thr Arg Pro Ala Pro Ala Ser Arg Val Ala

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20

Arg Met Ala Ser Asp Glu Phe Pro Ser Met Cys

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